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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/000,004A

DATE: 07/18/2001 TIME: 12:01:49

Input Set : A:\600-314SeqLst.txt

Output Set: N:\CRF3\07182001\I000004A.raw ENTERE

- 3 <110> APPLICANT: Tsilibary, Photini-Effie Charonis, Aristidis S. 5 Setty, Suman Mauer, Michael
- 8 <120> TITLE OF INVENTION: ANALYSIS OF ALPHA INTEGRINS FOR THE DIAGNOSIS OF DIABETIC NEPHROPATHY
 - 10 <130> FILE REFERENCE: 600.314USWO
 - 12 <140> CURRENT APPLICATION NUMBER: US 09/000,004A
- C--> 13 <141> CURRENT FILING DATE: 2001-06-19
 - 15 <150> PRIOR APPLICATION NUMBER: US 60/001,387
 - 16 <151> PRIOR FILING DATE: 1995-07-21
 - 18 <150> PRIOR APPLICATION NUMBER: US 60/001,861
 - 19 <151> PRIOR FILING DATE: 1995-08-03
 - 21 <150> PRIOR APPLICATION NUMBER: US 60/016,700
 - 22 <151> PRIOR FILING DATE: 1996-05-02
 - 24 <150> PRIOR APPLICATION NUMBER: PCT/US96/12067
 - 25 <151> PRIOR FILING DATE: 1996-07-19
 - 27 <160> NUMBER OF SEQ ID NOS: 16
 - 29 <170> SOFTWARE: PatentIn version 3.1
 - 31 <210> SEO ID NO: 1
 - 32 <211> LENGTH: 3989
 - 33 <212> TYPE: DNA
 - 34 <213> ORGANISM: Rattus
 - 36 <220> FEATURE:
 - 37 <221> NAME/KEY: CDS
 - 38 <222> LOCATION: (420)..(3959)
 - 39 <223> OTHER INFORMATION:
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73 ggt tot cot tta gtt ggc caa coc aaa gca aga act gga gat gtc tat

659



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74 75		Ser	Pro	Leu	Val	Gly 70	Gln	Pro	Lys	Ala	Arg 75	Thr	Gly	Asp	Val	Tyr 80	
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																	707
	гля	Cys	PIO	Val		AIG	GIU	ALG	Ата		PIO	Cys	val	пуъ	Leu 95	ASP	
79					85					90							755
	_														gaa		755
	Leu	Pro	Val		Thr	ser	тте	Pro		vaı	Thr	GIU	тте	_	Glu	Asn	
83	_			100					105					110			000
															ttt		803
	Met	Thr		GLY	Ser	Thr	Leu		Thr	Asn	Pro	Asn		GLY	Phe	Leu	•
87			115					120					125				
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90	Ala	-	Gly	Pro	Leu	\mathtt{Tyr}		Tyr	Arg	Cys	Gly		Leu	His	Tyr	Thr	
91		130					135					140					
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94	Thr	Gly	Ile	Cys	Ser	Asp	Val	Ser	Pro	Thr	Phe	Gln	Val	Val	Asn	Ser	
95	145					150					155					160	
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98	Phe	Ala	Pro	Val	Gln	Glu	Cys	Ser	Thr	Gln	Leu	Asp	Ile	Val	Ile	Val	
99					165					170					175		
101	. ctg	gat	ggc	tco	aac	ago	ato	: tac	ccc	t tgg	g gaa	ı agt	gto	ato	c gcc	ttt	995
102	Leu	Asp	Gly	Ser	Asr	Ser	: Ile	туз	r	Tr	Glu	ı Ser	· Val	l Ile	e Ala	Phe	
103	3			180)				185	5				190)		
105	tta	aac	gao	ctt	ctt	aag	agg	ato	g gat	att	ggc	cct	aag	cag	g aca	cag	1043
																Gln	
107			195			-	-	200	_		_	•	205				
109	gto	qqq	att	qta	cad	tat	. qqa	gad	, aat	gta	acc	cat	gad	tto	aac	ctc	1091
	_			_	_					_						Leu	
111		210				•	215					220					
113	aat	aaq	tat	tca	tco	aca	gaa	gad	qto	ctt	gto	gca	qca	aad	c aaa	ata	1139
																Ile	
	225	_	-			230					235				-	240	
			caq	aaa	aac	cto	caa	acc	rato	aca	ı qec	ctt	. qqa	ata	a qac	aca	1187
																Thr	
119	_			- 1	245					250			_		255		
		ago	aaa	σac			act	. qaa	ı act	cqc	r aat	gec	ago	aqq	ı qqa	gtt	1235
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123		9	-1-	260				0_0	265		,1			270			
		aaa	ato			att	ata	acc			σаа	t co	cat			tat	1283
															Asn		
127	_		275					280		, 011	010		285			1 -	
		cta			ato	ato	саа			σac	r dac	· αaa			cao	cga	1331
																Arg	1001
131		290		J_1.	. ,		295			. 510		300				3	
				act	ato	ctt			, tat	. 990	ann			, ++=	a a or c	act	1379
																Thr	13,3
	305		тте	ATO	. 116	310		HTS	, тут	. ASI	315	_	nol.	, ne	, Det	320	
			+++	a+ ~				227	+	+~			(T-2)		. 200	gaa	1427
																	142/
T 2 g	GIU	ьys	File	val	GIU	GIU	тте	- гу	ser	. тте	: нта	ser	GIU	LPIC	דווד י	Glu	



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139					325					330					335		
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143				340					345					350			
145	aaa	gct	ctg	gga	gaa	agg	ata	ttc	gct	ttg	gaa	gcg	aca	gct	gac	cag	1523
146	Lys	Ala	Leu	Gly	Glu	Arg	Ile	Phe	Ala	Leu	Glu	Ala	Thr	Ala	Asp	Gln	
147	_		355	_		_		360					365				
149	tca	qca	gct	tca	ttt	qaq	atq	qaa	atg	tct	caq	act	qqc	ttc	agt	gct	1571
														Phe			
151		370		-			375		-			380	_				
	cac		tac	cad	σac	taa	atc	atα	ctt	σσα	aca	ata	σσα	gcc	tat	gac	1619
														Ala			
	385	- y -	JCI	OLII	пор	390	vai	ricc	i.cu	O ₁	395	V 4 1	Gry	nra	- 1 -	400	
		220	~~~	act	ata		2+4	a a a	220	aat		G 2 G	2+4	gtc	ato		1667
														Val			1007
	пр	ASII	СТА	1111	405	Val	Mec	GTII	пуъ	410	ASII	GIII	мес	vai	415	PIO	
159																	1715
										-	_	_		gag		_	1715
	HIS	Asn	Thr		Pne	GIn	Thr	Glu		Ата	ьуs	Met	Asn	Glu	Pro	Leu	
163				420					425					430			
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	Ala	Ser	-	Leu	Gly	\mathtt{Tyr}	Thr		Asn	Ser	Ala	Thr		Pro	Gly	Asp	
167			435					440					445				
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170	Val	Leu	Tyr	Ile	Ala	Gly	Gln	Pro	Arg	Tyr	Asn	His	Thr	Gly	Gln	Val	
171		450					455					460					
173	gtc	atc	tac	aag	atg	gag	gat	ggg	aac	atc	aac	att	ctg	cag	aca	ctc	1859
174	Val	Ile	Tyr	Lys	Met	Glu	Asp	Gly	Asn	Ile	Asn	Ile	Leu	Gln	Thr	Leu	
175	465					470					475					480	
177	ggc	gga	gag	cag	att	ggt	tcc	tac	ttt	ggt	agt	gtc	tta	aca	aca	att	1907
178	Gly	Gly	Glu	Gln	Ile	Gly	Ser	Tyr	Phe	Gly	Ser	Val	Leu	Thr	Thr	Ile	
179	_	_			485	_		-		490					495		
181	qac	atc	gac	aaa	gat	tct	tat	act	gat	ctg	ctt	ctc	gtc	ggg	qcc	ccc	1955
182	Asp	Ile	Asp	Lys	Asp	Ser	Tyr	Thr	Asp	Leu	Leu	Leu	va1	Gly	Āla	Pro	
183	•		_	500	•		•		505					510			
	atσ	tac	ato	aaa	aca	gag	aaa	gag	gaa	caq	aac	aaσ	ata	tac	ata	tac	2003
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187	1100	-1-	515	U = 1		0	-10	520	0.2.0	V			525	-1-		-1-	
	act	ata		сал	aca	agg	+++	1.5	tat	caa	atα	acc		gaa	cca	att	2051
														Glu			2031
191	AIG	530	ASII	OIII	1111	2119	535	Olu	- 1 -	OIII	nec	540	пси	GIG	110	110	
	200		200	+ ~ ~	+ ~ ~	t a 2		ata	330	an+	22+		+ ~ ~	200	222	<i>α</i> 2 2	2099
		_		-	_					-				acg			2099
	_	GIII	TIIT	Cys	Cys		SET	Leu	гуѕ	ASP		ser	Cys	Thr	пÃР		
195			t	~		550	~~~	acc	~~~	++-	555	265	~	-++	~~+	560	2147
														att			2147
	ASN	гÀг	ASN	GIU		cys	дТΆ	нта	Arg		стλ	THE	АТА	Ile		AId	
199					565					570					575		0105
	_		_				-				-	-		att		-	2195
	val	rys	Asp		Asn	val	Asp	GLY		Asn	Asp	val	val	Ile	GLY	Ala	
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	ggc																2291
	GIA		Thr	He	Arg	GLu		Tyr	Ala	GIn	Arg		Pro	Ser	GLĀ	Gly	
211		610					615					620					
	gat																2339
	Asp	Gly	Lys	Thr	Leu		Phe	Phe	Gly	Gln		Ile	His	GTA	GLu		
	625					630					635					640	
	gat																2387
	Asp	Leu	Asn	Gly		Gly	Leu	Thr	Asp		Thr	Ile	Gly	GTA		GIY	
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	gga	_	-				_										2435
	Gly	Ala	Ala		Phe	\mathtt{Trp}	Ala	Arg		Val	Ala	Val	Val	_	Val	Thr	
223				660					665					670			
	atg			_													2483
226	Met	Asn	Phe	Glu	Pro	Asn	Lys	Val	Asn	Ile	Gln	Lys		Asn	Cys	Arg	
22.7			675					680					685				
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230	Val	Glu	Gly	Lys	Glu	Thr	Val	Cys	Ile	Asn	Ala	Thr	Met	Cys	Phe	His	
231	•	690					695					700					
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234	Val	Lys	Leu	Lys	Ser	Lys	Glu	Asp	Ser	Ile	Tyr	Glu	Ala	Asp	Leu	Gln	
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237	tac	cgt	gtc	acc	ctt	gat	tca	ctg	agg	cag	ata	tca	cgg	agc	ttt	ttt	2627
238	Tyr	Arg	Val	Thr	Leu	Asp	Ser	Leu	Arg	Gln	Ile	Ser	Arg	Ser	Phe	Phe	
239					725					730					735		
	tct																2675
242	Ser	Gly	Thr	Gln	Glu	Arg	Lys	Ile	Gln	Arg	Asn	Ile	Thr	Val	Arg	Glu	
243				740					745					750			
	tca	_	_														2723
246	Ser	Glu	Cys	Ile	Arg	His	Ser	Phe	Tyr	Met	Leu	Asp	Lys	His	Asp	Phe	
247			755					760					765				
	cag	-			_			-	_								2771
250	Gln	Asp	Ser	Val	Arg	Val		Leu	Asp	Phe	Asn		Thr	Asp	Pro	Glu	
251		770					775					780					
	aat																2819
254	Asn	Gly	Pro	Val	Leu	Asp	Asp	Ala	Leu	Pro	Asn	Ser	Val	His	Glu	His	
255						790					795					800	
	att																2867
258	Ile	${\tt Pro}$	Phe	Ala	Lys	Asp	Cys	Gly	Asn	Lys	Glu	Arg	Cys	Ile	Ser	Asp	
259					805					810			•		815		
	ctc																2915
262	Leu	Thr	Leu	Asn	Val	Ser	Thr	Thr	Glu	Lys	Ser	Leu	Leu	Ile	Val	Lys	
263				820		•			825					830			
	tcc																2963
266	Ser	Gln	His	Asp	Lys	Phe	Asn	Val	Ser	Leu	Thr	Val	Lys	Asn	Lys	Gly	
267			835					840					845				
269	gac	agt	gcg	tac	aac	acc	agg	aca	gtg	gtg	cag	cat	tca	cca	aat	ctg	3011

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	att		tca	gga	att	gag		atc	caa	aaa	qat			qaa	tct	aat	3059
																Asn	
	865			1		870				_1 -	875		- 1			880	
		aat.	atc	act.	tac		at.t.	σσα	tat	cct	t.t.c	cta	aσa	σca	σσα	gaa	3107
					_	_	_						-			Glu	
279	0.111	11011			885	5		1	-1-	890			,		895		
	acα	att	acc	ttc		ata	a t.a	t.t.c	caq		aac	aca	tcc	cat		tcg	3155
																Ser	
283				900	-1-				905			-		910			
	gaa	aat	qca	atc	att	cac	tta	aqt	qca	aca	agt	qac	agt	gag	qaq	ccc	3203
	-		-					_	_		-		-			Pro	
287			915					920				-	925				
	cta	gaa		ctt	aat	qat	aat	qaa	qta	aat	att	tcc	atc	cca	gta	aaa	3251
																Lys	
291		930				•	935					940				-	
	tat		att	qqa	ctq	caq	ttt	tac	agt	tct	qcq	agt	qaa	cat	cac	att	3299
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	945					950		_			955					960	
297	tca	qtc	qct	qcc	aat	gag	acq	atc	cct	gag	ttt	att	aac	tcc	act	gag	3347
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299					965					970					975		
301	qac	att	ggg	aat	gaa	att	aat	qtc	ttc	tat	acq	att	aga	aaq	agg	ggg	3395
	Asp				-			_			_		-	_			
	-									-							
303				980					985					990	_	-	
	cat	ttc	cca		cca	gaa	ctt	cag		j tca	a ati	t to		990		_	3443
305				atg					cto				a tt	990 c c	cc a	at ttg	3443
305				atg					cto Let				a tt	990 c c e P	cc a	at ttg	3443
305 306 307		Phe	Pro 995	atg Met	Pro	Glu	Leu	Gln 1000	cto Leo	ı Sei	r Ile	e Se	a tt r Ph	990 c c e P 05	cc a	at ttg sn Leu	3443 3488
305 306 307 309	His	Phe gca	Pro 995 gat	atg Met	Pro tat	Glu cct	Leu . gta	Gln 1000	cto Leo) ig ta	sei	r Ile	e Se tt g	a tter Phe 10	990 c c e P 05 tgg	cc a ro A	at ttg sn Leu tct	
305 306 307 309	His	Phe gca	Pro 995 gat Asp	atg Met	Pro tat	Glu cct	Leu . gta	Gln 1000 ct Le	cto Leo) ig ta	sei	r Ile	e Se tt g le G	a tter Phe 10	990 c c e P 05 tgg	cc a ro A tca	at ttg sn Leu tct	
305 306 307 309 310 311	His	Phe gca Ala 1010	Pro 995 gat Asp	atg Met ggt	Pro tat Tyr	Glu cct Pro	Leu gta Val	Gln 1000 a ct L Le	cto Leo) tg ta eu Ty	ı Sen ac co yr Pi	r Ile ca at ro Il	e Se tt g le G 1	a tt r Pho 10 ga ly 6	990 c c e P 05 tgg Trp	cc a ro A tca	at ttg sn Leu tct Ser	
305 306 307 309 310 311 313	His acg Thr	Phe gca Ala 1010 gat	Pro 995 gat Asp) aat	atg Met ggt Gly	Pro tat Tyr	Glu cct Pro	Leu gta Val 101 aga	Gln 1000 L Ct L6	cto Let) ig to eu Ty	ser ac co yr Pr gg ag	r Ile ca at ro Il	e Se tt g le G 1 tt g	a tter Pho 10 ga ly 6 020	990 c c e P 05 tgg Trp	cc a ro A tca Ser	at ttg sn Leu tct Ser	3488
305 306 307 309 310 311 313	His acg Thr tca	Phe gca Ala 1010 gat	Pro 995 gat Asp aat Asr	atg Met ggt Gly	Pro tat Tyr	Glu cct Pro	Leu gta Val 101 aga	Gln 1000 L Ct L6	cto Let) ig to eu Ty	ser ac co yr Pr gg ag	r Ile ca at ro Il	e Se tt g le G tt g eu G	a tter Pho 10 ga ly 6 020	990 c c e P 05 tgg Trp	cc a ro A tca Ser	at ttg sn Leu tct Ser	3488
305 306 307 309 310 311 313 314 315	His acg Thr tca	gca Ala 1010 gat Asp 1025	Pro 995 gat Asp aat Asr	atg Met ggt Gly	Pro tat Tyr aac Asr	Glu cct Pro tgt	Leu gta Val 101 aga Arg	Gln 1000 1 Ct L Le 5 C0 1 P1	cto Let Let Cc co Cc co Co Ai	ac co yr Pi gg ag	r Ile ca at ro I. gc ct er Le	e Se tt g le G tt g eu G	a ttr r Pho 10 ga ly '020 ag '035 ag '035	990 c c e P 05 tgg Trp gac Asp	cc a ro A tca Ser ccc Pro	at ttg sn Leu tct Ser ttt Phe	3488
305 306 307 309 310 311 313 314 315 317	His acg Thr tca Ser	gca Ala 1010 gat Asp 1025 atc	Pro 995 gat Asp aat Asr	atg Met ggt Gly gtg Val	Pro tat Tyr aac Asr	Glu cct Pro tgt Cys	Leu 2 gta 3 Val 3 101 4 aga 4 Arg 4 103	Gln 1000 a ct Le 5 a cc a cc y P1 80	cto Led)) eu Ty cc co co Al	i Sei ac co yr Pi gg ag cg Se	r Ileca at ro II	tt golden from the second seco	a tt r Pho 10 ga ly 020 ag lu 035 ag	990 c c e P 05 tgg Trp gac Asp	cc a ro A tca Ser ccc Pro	at ttg sn Leu tct Ser ttt Phe	3488 3533
305 306 307 309 310 311 313 314 315 317	His acg Thr tca Ser	gca Ala 1010 gat Asp 1025 atc	Pro 995 gat Asp aat Asn aac	atg Met ggt Gly gtg Val	Pro tat Tyr aac Asr	Glu cct Pro tgt Cys	Leu 2 gta 3 Val 3 101 4 aga 4 Arg 4 103	Gln 1000 1 ct Le 5 1 cc 1 pr 1000 1 at 1000 1 at 1000 1000 1000 1000 1000 1000 1000 10	cto Lev) ccg ta eu Ty ccc co cc co	i Sei ac co yr Pi gg ag cg Se	r Ileca at ro II	tt gle G tt ge tt ge tt g tt g tt g tt g tt g tt	a tt r Pho 10 ga ly 020 ag lu 035 ag	990 c c e P 05 tgg Trp gac Asp	cc a ro A tca Ser ccc Pro	at ttg sn Leu tct Ser ttt Phe	3488 3533
305 306 307 309 310 311 313 314 315 317 318 319	His acg Thr tca Ser	gca Ala 1010 gat Asp 1025 atc Ile 1040	Pro 995 gat Asp aat Asr aac Asr	atg Met ggt Gly Val tct Ser	Pro tat Tyr aac Asr	Glu cct Pro tgt Cys aag	Leu gta Val 101 aga Arg 103 aaa Lys 104 cag	Gln 1000 1 ct L6 5 1 cc y Pi 80 1 at 8 Me 15	cto Let Cto Cto Cto Cto Cto Cto Cto Cto Cto C	n Sen ac co yr Pr gg ag rg Se ca at hr II	r Ileca at control II	tt g le G tt g eu G cg a er L gt a	a ttr r Pho ga ly 020 ag 035 ag ys 050 cg	990 c c e P 05 tgg Trp gac Asp	cc a ro A tca Ser ccc Pro	at ttg sn Leu tct Ser ttt Phe gtt Val	3488 3533
305 306 307 309 310 311 313 314 315 317 318 319 321	acg Thr tca Ser ggc Gly	gca Ala 1010 gat Asp 1025 atc Ile 1040 aaa	Pro 995 gat Asp aat Asr aac Asr	atg Met ggt Gly Val tct Ser	Pro tat Tyr aac Asr	Glu cet Pro tgt Cys aag	Leu gta Val 101 aga Arg 103 aaa Lys 104 cag	Gln 1000 1 ct L6 5 1 cc y Pi 80 1 at 8 Me 15	cto Let Cto Cto Cto Cto Cto Cto Cto Cto Cto C	n Sen ac co yr Pr gg ag rg Se ca at hr II	r Ileca at control of the control of	tt g le G tt g eu G cg a er L gt a	a ttr r Pho ga ly 020 ag 035 ag ys 050 cg	990 c c e P 05 tgg Trp gac Asp tct Ser	cc a ro A tca Ser ccc Pro gag Glu	at ttg sn Leu tct Ser ttt Phe gtt Val	3488 3533 3578
305 306 307 309 310 311 313 314 315 317 318 319 321	acg Thr tca Ser ggc Gly	gca Ala 1010 gat Asp 1025 atc Ile 1040 aaa	Pro 995 gat Asp aat Asr aac Asr aac Asr	atg Met ggt Gly Val tct Ser	Pro tat Tyr aac Asr	Glu cct Pro tgt Cys aag	Leu gta Val 101 aga Arg 103 aaa Lys 104 cag	Gln 1000 a ct Le 5 a cc B0 a t S B0 a t S Me 15	cto Let Cto Cto Cto Cto Cto Cto Cto Cto Cto C	n Sen ac co yr Pr gg ag rg Se ca at hr II	r Ileca at control II	tt g le G tt g eu G cg a er L gt a gt a	a ttr r Pho ga ly 020 ag 035 ag ys 050 cg	990 c c e P 05 tgg Trp gac Asp tct Ser	cc a ro A tca ser ccc Pro gag Glu gga	at ttg sn Leu tct Ser ttt Phe gtt Val	3488 3533 3578 3623
305 306 307 309 310 311 313 314 315 317 318 321 322 323 325	acg Thr tca Ser ggc Gly ctc Leu	gca Ala 1010 gat Asp 1025 atc Ile 1040 aaa Lys acc	Pro 995 gat Asp aat Asn aac Asn aga Arg	atg Met ggt Gly Val ctct Ser ggc	tat Tyr aac Asr Gly	Glu cct Pro tgt Cys aag	Leu gta Val 101 aga Arg 103 aaa Lys 104 cage Glr 106	Gln 1000 1 Ct L6 5 1 C0 80 1 at 8 Me 15 9 ga 1 As	cto Let) cg ta eu Ty cc co co Al cg ac et Th ac to	ac coryr Program of Services o	r Ileca at control II	tt g le G tt g eu G cg a er L gt a gt a er 1	a ttr Phoga 10 20 ag 10 35 ag ys 5050 cg hr 6065 tg	990 c c e P 05 tgg Trp gac Asp tct Ser tgt	cc a ro A tca ser ecc Pro gag Glu gga Gly caa	at ttg sn Leu tct Ser ttt Phe gtt Val gtt Val	3488 3533 3578
305 306 307 309 310 311 313 314 315 317 318 321 322 323 325 326	acg Thr tca Ser ggc Gly ctc Leu	gca Ala 1010 gat Asp 1025 atc Ile 1040 aaa Lys acc Thr	Pro 995 gat Asp aat Asn aac Asn aga Arg	atg Met ggt ggt Val tct Ser ggc ggc	Pro tat Tyr aac Asr ggg Gly aca Thr	Glu cet pro ctgt Cys aag Lys atc	Leu gta Val 101 aga Arg 103 aaa Lys 104 cag Glr 106	Gln 1000 1 Ct L6 5 1 Cc 15 1 Cc 15 1 G 15 1 As 15 1 G 16 1 L 16 16 17 17 18 18 18 18 18 18 18 18 18 18 18 18 18	cto Let) cg ta eu Ty cc co cro Ai cc to et Th ac to	ac coryr Program ac coryr Program ac cory Seca at ac corys Seca at ac corys Seca ac corps Seca ac corys Seca ac corps Seca ac corys Seca ac corps Seca ac corys Seca ac co	ca at collect the Second can be second can b	tt g le G tt g eu G er L g g t a g er L g tc C g tc C sp L	a ttr Phoga 10 20 ag 10 35 ag 950 cg hr 0065 tg 6eu	990 c c e P 05 tgg Trp gac Asp tct Ser tgt	cc a ro A tca Ser ccc Pro gag Glu gga Gly	at ttg sn Leu tct Ser ttt Phe gtt Val gtt Val	3488 3533 3578 3623
305 306 307 309 310 311 313 314 315 317 318 319 321 322 323 325 326 327	acg Thr tca Ser ggc Gly ctc Leu gcc Ala	gca Ala 1010 gat Asp 1025 atc Ile 1040 aaa Lys acc Thr 1070	Pro 995 gat Asr aat Asr aac Asr aga arg	atg Met ggt ggt Val tct Ser ggc Gly acg	tat Tyr aac Asr Gly aca Thr	cet cet cet cet cet cet cet cet cet cet	Leu gta yal 101 aga Arg 103 aaa Lys 104 cag Glr 106	Gln 1000 1 Ct L6 5 1 Co 1 P1 80 1 at 5 Me 15 1 G 1 Ct 1 Le 17 1 Ct	cto Let Cg ta cg ta cc co cc A cg ac cd to c	ac coyr Progg acg set to set to	ca at correct second se	tt g	a ttr Phoga 19 020 ag 10 035 ag 9050 cg hr 0065 tg 6080	990 c c e P 05 tgg Trp gac Asp tct Ser tgt	cc a ro A tca Ser ccc Pro gag Glu gga Gly caa Gln	at ttg sn Leu tct Ser ttt Phe gtt Val gtt Val gtg Val	3488 3533 3578 3623 3668
305 306 307 309 310 311 313 314 315 317 318 319 321 322 323 325 326 327 329	acg Thr tca ser ggc Gly ctc Leu gcc Ala	gca Ala 1010 gat Asp 1025 atc Ile 1040 aaa Lys 1055 acc Thr 1070 gtc	Pro 995 gat Asr aat Asr aac Asr aga Arg tcg	atg Met ggt ggt Val tct Ser ggc Gly acg	tat Tyr aac Asr Gly aca Thr	ctgt cys aag Lys atc ser ctg	gta yal 101 aga Arg 103 aaa Lys 104 cag Ctc 107	Gln 1000 1 Ct Le 5 1 Co 1 P1 80 1 at 5 1 Ga 15 1 Ga 15 1 Le 17 1 at 17	cto Let Cg ta cc co cc co cc co cc to	ac corrections of the correction	ca at ro II gc ct gc ct ta to le Se gt age cc ga cc ga cc tt	tt gg tt gg tu G er L gt a er T gt a er T ac C ac p L ac c ac c ac a	a ttr Phoga 19 020 ag 10 35 ag 950 cg hr 065 tg 680 ta 6	990 c c e P 05 trp gac Asp tct Ser tgt cys	cc a ro A tca ser ccc Pro gag Glu gga Gly caa Gln gca	at ttg sn Leu tct Ser ttt Phe gtt Val gtt Val gtg Val cat	3488 3533 3578 3623
305 306 307 310 311 313 314 315 317 318 319 321 322 323 325 326 327 329 330	acg Thr tca Ser ggc Gly ctc Leu gcc Ala	gca Ala 1010 gat Asp 1025 atc Ile 1040 aaa Lys 1055 acc Thr 1070 gtc Val	Pro 995 gat Asr aat Asr aga Arg atc Ile	atg Met ggt ggt Val tct Ser ggc Gly acg	tat Tyr aac Asr Gly aca Thr	cet cet cet cet cet cet cet cet cet cet	Leu gta yal 101 aga Arg 103 aaa Lys 104 cag Ctc 107 tgg	Gln 1000 1 Ct LE 5 1 Co 130 1 at 5 1 G 15 1 G 15 1 G 15 1 L 15 1 G 15 1 L 1 L 1 L 1 L 1 L 1 L 1 L 1 L 1 L 1 L	cto Let Cg ta cc co cc co cc co cc to	ac corrections of the correction	ca at correct second se	tt gg tt gg te G tt gg te G tt gg te T gt ac er T ac C ac T ac C ac T ac C ac T	a ttr Phoga 19 19 10 20 ag 10 35 ag 10 50 cg hr 60 65 tg eu 10 80 ta 10 10 10 10 10 10 10 10 10 10 10 10 10	990 c c e P 05 trp gac Asp tct Ser tgt cys	cc a ro A tca Ser ccc Pro gag Glu gga Gly caa Gln	at ttg sn Leu tct Ser ttt Phe gtt Val gtt Val gtg Val cat	3488 3533 3578 3623 3668
305 306 307 310 311 313 314 315 317 318 319 321 322 323 325 326 327 329 330 331	His acg Thr tca ser ggc Gly ctc Leu gcc Ala aat Asn	gca Ala 1010 gat Asp 1025 atc Ile 1040 aaa Lys 1055 acc Thr 1070 gtc Val 1085	Pro 995 gat Asr aat Asr aga Arg atc Ile	atg Met ggt Gly Val tct Gly Gly Cacg Thr	tat Tyr aac Asr Gly aca Thr tgt	ctgt cys aag Lys atc ctg	Leu gta lota lota lota lota lota lota lota lo	Gln 1000 1 Ct L Le 5 1 Co 1 P1 100 1 at 15 1 Ga 1 As 15 1 Ga 1 Le 15 1 Le 16 16 16 16 16 16 16 16 16 16 16 16 16	cto Let Cg ta cg ta cc co co Al cg ac co Al cc to	ac corrections of the correction	ca at the Secretary Secret	tt gg tt gg teg 1 tt gg teg 1 gt ac er 1 ac cc ac fi tc ac ine I:	a ttr Pho 10 ga 1y 020 ag 10 35 ag 950 cg hr 065 tg 6080 ta 20 10 5	990 c c e P 05 trp gac Asp tct Ser tys agt	cc a ro A tca Ser ccc Pro gag Glu gga Gly caa Gln gca Ala	at ttg sn Leu tct Ser ttt Phe gtt Val gtt Val gtg Val cat His	3488 3533 3578 3623 3668 3713
305 306 307 310 311 313 314 315 317 318 319 321 325 326 327 329 330 331 333	acg Thr tca ser ggc Gly ctc Leu gcc Ala	gca Ala 1010 gat Asp 1025 atc Ile 1040 aaa Lys 1055 acc Thr 1070 gtc Val 1085	Pro 995 gat Asr aat Asr aac Asr i aga Arg i tcg ser	atg Met ggt Gly Lgtg Val Lstct Gly Lacg Thr Leu Lta	tat Tyr aac Asr Gly aca Thr tgt Cys	ctgt cys aag Lys atc ser ctg	Leu gta loga 101 aga Arg 103 aaa Lys 104 cag ctc 107 tgg Trp 109 act	Gln 1000 1 Ct L Le 5 1 Cc J P1 80 1 at 5 J Ga 60 C Ct J Ct	cto Let Cec co Co Co Al Cec co	ac corrections of the correction	ca at ro II gc ct gc ct ta to le Se gt age cc ga cc ga cc tt	tt gg tt gg te G tt gg te G tt gg te T tt ac er 1 tc ac tc a	a ttr Phoga 19 10 20 ag 10 20	990 c c e P 05 trp gac Asp tct Ser tgt agt	cc a ro A tca ser ccc Pro gag Glu gga Gly caa Gln gca	at ttg sn Leu tct Ser ttt Phe gtt Val gtt Val gtg Val cat His	3488 3533 3578 3623 3668

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/000,004A

DATE: 07/18/2001

TIME: 12:01:50

Input Set : A:\600-314SeqLst.txt

Output Set: N:\CRF3\07182001\1000004A.raw

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date